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Full Record

Details for HUGENEFL:L17131_RNA1_AT

Full Screen

NetAffx Links	Cluster Members Consensus/Exemplar

GeneChip Array Information

Probe Set ID	L17131_rna1_at
GeneChip Array	HumanGeneFL Array
Organism Common Name	Human

Probe Design Information

Transcript ID	L17131_rna1
Sequence Type	Exemplar sequence
Representative Public ID	L17131 NCBI
Target Description	L17131, class A, 20 probes, 20 in L17131mRNA#1 1646-2198, Human high mobility group protein (HMG-I(Y)) gene exons 1-8, complete cds

Sequence

Target Sequence

```
>HUGENEFL:L17131_RNA1_AT
ttgtccaggtgaggcccaagagccctgtggccgccacctgaggtgggctggggctgctcc
cctaaccctactttcgttccgccactcagccatttccccctcctcagatggggcaccaat
aacaaggagctcaccctgcccgctcccaaccccctcctgctcctccctgccccccaagg
ttctggttccatttttcctctgttcacaaactacctctggacagttgtgttgttttttgt
tcaatgttccattcttcgacatccgtcattgctgctgctaccagcgccaaatgttcatcc
tcattgcctcctgttctgcccacgatcccctcccccaagatactctttgtggggaagagg
ggctggggcatggcaggctgggtgaccgactaccccagtcccagggaaggtggggccctg
cccctaggatgctgcagcagagtgagcaaggggcccgaatcgaccataaagggtgtagg
ggccacctcctcccccctgttctgttggggaggggtagccatgatttgtcccagcctgggg
ctccctctctggtttcctatttgcagttacttgaata
```

Probe Sequence(5'-3')	Probe X	Probe Y	Probe Interrogation Position	Strandedness
TTGTCCAGGTGAGGCCCAAGAGCCC	294	101	1658	Antisense
AGGTGAGGCCCAAGAGCCCTGTGGC	295	101	1664	Antisense
ACCAATAACAAGGAGCTCACCCTGC	296	101	1772	Antisense
TTTTCCTCTGTTCACAAACTACCTC	297	101	1850	Antisense
CTACCTCTGGACAGTTGTGTTGTTT	298	101	1868	Antisense
TTCCATTCTTCGACATCCGTCATTG	299	101	1904	Antisense
TCTTCGACATCCGTCATTGCTGCTG	300	101	1910	Antisense

	GCTACCAGCGCCAAATGTTCATCCT	301	101	1934	Antisense
	TCATCCTCATTGCCTCCTGTTCTGC	302	101	1952	Antisense
	TCATTGCCTCCTGTTCTGCCCACGA	303	101	1958	Antisense
	AAGATACTCTTTGTGGGGAAGAGGG	304	101	1994	Antisense
	GCAGGCTGGGTGACCGACTACCCCA	305	101	2030	Antisense
	CCCCTAGGATGCTGCAGCAGAGTGA	306	101	2078	Antisense
Probe Info	AGCAAGGGGGCCCGAATCGACCATA	307	101	2102	Antisense
	CGAATCGACCATAAAGGGTGTAGGG	308	101	2114	Antisense
	GCCATGATTTGTCCCAGCCTGGGGC	309	101	2174	Antisense
	CTGGGGCTCCCTCTCTGGTTTCCTA	310	101	2192	Antisense
	CTCCCTCTCTGGTTTCCTATTTGCA	311	101	2198	Antisense
	CTCTGGTTTCCTATTTGCAGTTACT	312	101	2204	Antisense
	TTTCCTATTTGCAGTTACTTGAATA	313	101	2210	Antisense